

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/830,502Source: 1600Date Processed by STIC: 4/24/2003

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry directly to:
 U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003



1600

RAW SEQUENCE LISTING DATE: 04/24/2003 PATENT APPLICATION: US/09/830,502 TIME: 14:25:05

Input Set: A:\C26151.app

Output Set: N:\CRF4\04242003\1830502.raw

- 3 <110> APPLICANT: Barany, Francis Cao, Weiguo Tong, Jie
- 7 <120> TITLE OF INVENTION: HIGH FIDELITY THERMOSTABLE LIGASE AND USES THEREOF
- 9 <130> FILE REFERENCE: 19603/2615
- 11 <140> CURRENT APPLICATION NUMBER: 09/830,502
- 12 <141> CURRENT FILING DATE: 1999-10-29
- 14 <150> PRIOR APPLICATION NUMBER: 60/106,461
- 15 <151> PRIOR FILING DATE: 1998-10-30
- 17 <150> PRIOR APPLICATION NUMBER: PCT/US99/25437
- 18 <151> PRIOR FILING DATE: 1999-10-29
- 20 <160> NUMBER OF SEQ ID NOS: 20
- 22 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

| | | | | _ | D NO H: 18 | : 15 84 | | | | | | Does Not Comply Corrected Diskette Needed | | | | | | |
|----|-----|------|-------|-------|---------------|------------|------|------|------|-----|-----|---|-----|-----|-----|------|-----|---------|
| | 453 | <21 | 2> T | YPE: | PRT | | | | | | | | | _ | 1 | 11 (| 5-7 | |
| | 454 | <21 | 3> 01 | RGAN: | ISM: | The: | rmus | aqua | atic | us | | | 1 | pp | 1- | T, 4 |) / | |
| | 456 | <400 | 0> SI | EQUE | NCE: | 15 | | - | | | | | - 0 | , , | | • | | |
| • | | | | | | | Lvs | Val | asA | Gly | Leu | Ser | Val | Asn | Leu | Tvr | Tvr | |
| | 458 | 1 | | | | 5 | _ | | - | - 2 | 10 | | | | | 15 | 4 | |
| E> | | Glu | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | · × |
| | 461 | | | | 20 | | | | | 25 | | | | | 30 | | | |
| E> | | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | |
| | 464 | | | 35 | | | | | 40 | | | | | 45 | | | | |
| E> | 466 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | |
| | 467 | | 50 | | | | | 55 | | | | | 60 | | | | | 0 |
| E> | 469 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Λe |
| | 470 | 65 | | | | | 70 | | | | | 75 | | | | | 80 | 10 |
| E> | 472 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | |
| | 473 | | | | | 85 | | | | | 90 | | | | | 95 | | \mathre |
| E> | 475 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | |
| | 476 | | | | 100 | | | | | 105 | | | | | 110 | | | |
| E> | 478 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Leu | Glu | Glu | Thr | Glv | Xaa | Xaa | Xaa | |
| | 479 | | | 115 | | | | | 120 | | | | | 125 | | | | |
| E> | | Xaa | Xaa | | Xaa | Xaa | Xaa | Xaa | | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | |
| | 482 | | 130 | | | | | 135 | | | | | 140 | | | | | |
| E> | 484 | Xaa | | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | |
| - | 485 | | | | | | 150 | | | | | 155 | | | | | 160 | |
| E> | | | Xaa | Xaa | Xaa | Xaa | | Xaa | Xaa | Xaa | Yaa | | Xaa | Pro | Phe | Glu | | |
| _ | | | | | | | | | | | | | | | | 4 | | |

Laa's reed to be explained. See p. 6 for more information

Does Not Comply

RAW SEQUENCE LISTING DATE: 04/24/2003 PATENT APPLICATION: US/09/830,502 TIME: 14:25:05

Input Set: A:\C26151.app
Output Set: N:\CRF4\04242003\1830502.raw

488 175 165 170 490 Asp Gly Val Val Lys Leu Asp 180 494 <210> SEQ ID NO: 16 495 <211> LENGTH: 187 496 <212> TYPE: PRT 497 <213> ORGANISM: Thermus flavus 499 <400> SEQUENCE: 16 500 Tyr Thr Val Glu His Lys Val Asp Gly Leu Ser Val Asn Leu Tyr Tyr 501 1 20 25 55 513 65 E--> 521 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Leu Glu Glu Val Glu Arg Glu Gly 115 120 135 140 528 145 150 155 165 533 Phe Glu Ala Asp Gly Val Val Lys Leu Asp 534 180 537 <210> SEQ ID NO: 17 538 <211> LENGTH: 184 539 <212> TYPE: PRT 540 <213> ORGANISM: Thermus filiformis 542 <400> SEQUENCE: 17 543 Tyr Thr Val Glu His Lys Val Asp Gly Leu Ser Val Asn Leu Tyr Tyr 544 1 10 same eno 35 40 45 556 65 70

per P.3

85

559

RAW SEQUENCE LISTING DATE: 04/24/2003 PATENT APPLICATION: US/09/830,502 TIME: 14:25:06

Input Set: A:\C26151.app

Output Set: N:\CRF4\04242003\I830502.raw

100 105 E--> 564 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Leu Glu Glu Ser Gly Xaa Xaa Xaa 115 120 125 565 135 571 145 150 155 165 170 576 Asp Gly Val Val Lys Met Asp 577 180 580 <210> SEO ID NO: 18 581 <211> LENGTH: 184 582 <212> TYPE: PRT 583 <213> ORGANISM: Thermus filiformis 585 <400> SEQUENCE: 18 586 Tyr Thr Val Glu His Lys Val Asp Gly Leu Ser Val Asn Leu Tyr Tyr 587 5 10 20 25 593 35 40 596 50 55 599 65 70 75 85 90 605 100 105 E--> 607 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Leu Glu Ser Gly Xaa Xaa Xaa 608 115 120 125 135 140 150 155 165 170 619 Asp Gly Val Val Val Lys Leu Asp 620 . 180 623 <210> SEQ ID NO: 19 624 <211> LENGTH: 184 625 <212> TYPE: PRT 626 <213> ORGANISM: Thermus sp. 628 <400> SEQUENCE: 19 629 Tyr Thr Val Glu His Lys Val Asp Gly Leu Ser Val Asn Leu Tyr Tyr 10

Same

p.4

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/830,502

DATE: 04/24/2003 TIME: 14:25:06

Input Set : A:\C26151.app

Output Set: N:\CRF4\04242003\1830502.raw

| | 633 | | | | 20 | | | | | 25 | | | | | 30 | | |
|-------------------|--|---|---|---|---|--|--|--|--|--|--|---|---|--|--|--|--|
| E> | 635 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa |
| | 636 | | | 35 | | | | | 40 | | | | | 45 | | | |
| E> | 638 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa |
| | 639 | | 50 | | | | | 55 | | | | | 60 | | | | |
| E> | 641 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa |
| | 642 | 65 | | | | | 70 | | | | | 75 | | | | | 80 |
| E> | 644 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa |
| | 645 | | | | | 85 | | | | | 90 | | | | | 95 | |
| E> | 647 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa |
| | 648 | | | | 100 | | | | | 105 | | | | | 110 | | |
| E> | 650 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Leu | Glu | Glu | Ser | Gly | Xaa | Xaa | Xaa |
| | 651 | | | 115 | | | | | 120 | | | | | 125 | | | |
| E> | 653 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa |
| | 654 | | 130 | | | | | 135 | | | | | 140 | | | | |
| E> | 656 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa |
| | 657 | 145 | | | | | 150 | | | * | | 155 | | | | | 160 |
| E> | 659 | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Pro | Phe | Glu | Ala |
| | 660 | | | | | 165 | | | | | 170 | | | | | 175 | |
| | 662 | Asp | Gly | Val | Val | Val | Lys | Leu | Asp | | | | | | | | |
| | 663 | - | - | | 180 | | - | | - | | | | | | | | |
| | 666 | <210 |)> SI | EQ II | ON C | : 20 | | | | | | | | | | | |
| | 667 | <213 | L> LE | ENGT | 1: 18 | 34 | | | | | | | | | | | |
| | 668 | <212 | 2> T | YPE: | PRT | | | | | | | | | | | | |
| | | | | RGAN | | The | cmus | sp. | | | | | | | | | |
| | | | | EQUE | | | | - | | | | | | | | | |
| | ~ - ~ | | | | | | | | | | _ | | | | | | |
| | 672 | Tyr | Thr | Val | Glu | His | Lys | Val | Asp | Gly | Leu | Ser | Val | Asn | Leu | Tyr | Tyr |
| | 673 | Tyr 1 | Thr | Val | Glu | His 5 | Lys | Val | Asp | Gly | Leu 10 | Ser | Val | Asn | Leu | Tyr 15 | Tyr |
| E> | 673 | 1 | | | | 5 | _ | | _ | | 10 | | | | | 15 | |
| E> | 673 | 1 | | | | 5 | _ | | _ | | 10 | | | | | 15 | |
| E> | 673 675 676 | 1 Glu | Xaa | Xaa | Xaa 20 | 5 Xaa | Xaa | Xaa | Xaa | Xaa 25 | 10 Xaa | Xaa | Xaa | Xaa | Xaa 30 | 15 Xaa | Xaa |
| | 673 675 676 | 1 Glu | Xaa | Xaa | Xaa 20 | 5 Xaa | Xaa | Xaa | Xaa | Xaa 25 | 10 Xaa | Xaa | Xaa | Xaa | Xaa 30 | 15 Xaa | Xaa |
| | 673 675 676 678 679 | 1 Glu Xaa | Xaa Xaa | Xaa Xaa 35 | Xaa 20 Xaa | 5 Xaa Xaa | Xaa Xaa | Xaa Xaa | Xaa Xaa 40 | Xaa 25 Xaa | 10 Xaa Xaa | Xaa Xaa | Xaa Xaa | Xaa Xaa 45 | Xaa 30 Xaa | 15 Xaa Xaa | Xaa Xaa |
| E> | 673 675 676 678 679 | 1 Glu Xaa | Xaa Xaa | Xaa Xaa 35 | Xaa 20 Xaa | 5 Xaa Xaa | Xaa Xaa | Xaa Xaa | Xaa Xaa 40 | Xaa 25 Xaa | 10 Xaa Xaa | Xaa Xaa | Xaa Xaa | Xaa Xaa 45 | Xaa 30 Xaa | 15 Xaa Xaa | Xaa Xaa |
| E> | 673 675 676 678 679 681 682 | 1 Glu Xaa Xaa | Xaa Xaa Xaa 50 | Xaa Xaa 35 Xaa | Xaa 20 Xaa Xaa | 5 Xaa Xaa Xaa | Xaa Xaa Xaa | Xaa Xaa Xaa 55 | Xaa Xaa 40 Xaa | Xaa 25 Xaa Xaa | 10 Xaa Xaa Xaa | Xaa Xaa Xaa | Xaa Xaa Xaa 60 | Xaa Xaa 45 Xaa | Xaa 30 Xaa Xaa | 15 Xaa Xaa Xaa | Xaa Xaa Xaa |
| E> | 673 675 676 678 679 681 682 | 1 Glu Xaa Xaa | Xaa Xaa Xaa 50 | Xaa Xaa 35 Xaa | Xaa 20 Xaa Xaa | 5 Xaa Xaa Xaa | Xaa Xaa Xaa | Xaa Xaa Xaa 55 | Xaa Xaa 40 Xaa | Xaa 25 Xaa Xaa | 10 Xaa Xaa Xaa | Xaa Xaa Xaa | Xaa Xaa Xaa 60 | Xaa Xaa 45 Xaa | Xaa 30 Xaa Xaa | 15 Xaa Xaa Xaa | Xaa Xaa Xaa |
| E> | 673 675 676 678 679 681 682 684 685 | 1 Glu Xaa Xaa Xaa 65 | Xaa Xaa Xaa 50 Xaa | Xaa Xaa 35 Xaa Xaa | Xaa 20 Xaa Xaa Xaa | 5 Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa 70 | Xaa Xaa Xaa 55 Xaa | Xaa Xaa 40 Xaa Xaa | Xaa 25 Xaa Xaa Xaa | 10 Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa 75 | Xaa Xaa Xaa 60 Xaa | Xaa Xaa 45 Xaa Xaa | Xaa 30 Xaa Xaa Xaa | 15 Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa 80 |
| E> E> | 673 675 676 678 679 681 682 684 685 | 1 Glu Xaa Xaa Xaa 65 | Xaa Xaa Xaa 50 Xaa | Xaa Xaa 35 Xaa Xaa | Xaa 20 Xaa Xaa Xaa | 5 Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa 70 | Xaa Xaa Xaa 55 Xaa | Xaa Xaa 40 Xaa Xaa | Xaa 25 Xaa Xaa Xaa | 10 Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa 75 | Xaa Xaa Xaa 60 Xaa | Xaa Xaa 45 Xaa Xaa | Xaa 30 Xaa Xaa Xaa | 15 Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa 80 |
| E> E> | 673 675 676 678 679 681 682 684 685 | 1 Glu Xaa Xaa Xaa 65 Xaa | Xaa Xaa So Xaa Xaa | Xaa Xaa 35 Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa | 5 Xaa Xaa Xaa Xaa Xaa 85 | Xaa Xaa Xaa Xaa 70 Xaa | Xaa Xaa Xaa 55 Xaa Xaa | Xaa Xaa 40 Xaa Xaa Xaa | Xaa 25 Xaa Xaa Xaa | 10 Xaa Xaa Xaa Xaa Xaa 90 | Xaa Xaa Xaa Xaa 75 Xaa | Xaa Xaa 60 Xaa Xaa | Xaa Xaa 45 Xaa Xaa Xaa | Xaa 30 Xaa Xaa Xaa Xaa | 15 Xaa Xaa Xaa Xaa 95 | Xaa Xaa Xaa 80 Xaa |
| E> E> E> | 673 675 676 678 679 681 682 684 685 | 1 Glu Xaa Xaa Xaa 65 Xaa | Xaa Xaa So Xaa Xaa | Xaa Xaa 35 Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa | 5 Xaa Xaa Xaa Xaa Xaa 85 | Xaa Xaa Xaa Xaa 70 Xaa | Xaa Xaa Xaa 55 Xaa Xaa | Xaa Xaa 40 Xaa Xaa Xaa | Xaa 25 Xaa Xaa Xaa | 10 Xaa Xaa Xaa Xaa Xaa 90 | Xaa Xaa Xaa Xaa 75 Xaa | Xaa Xaa 60 Xaa Xaa | Xaa Xaa 45 Xaa Xaa Xaa | Xaa 30 Xaa Xaa Xaa Xaa | 15 Xaa Xaa Xaa Xaa 95 | Xaa Xaa Xaa 80 Xaa |
| E> E> E> | 673 675 676 678 679 681 682 684 685 687 688 | 1 Glu Xaa Xaa Xaa 65 Xaa Xaa | Xaa Xaa 50 Xaa Xaa Xaa | Xaa Xaa 35 Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa Xaa 85 Xaa | Xaa Xaa Xaa 70 Xaa Xaa | Xaa Xaa 55 Xaa Xaa Xaa | Xaa Xaa 40 Xaa Xaa Xaa | Xaa 25 Xaa Xaa Xaa Xaa Xaa | 10 Xaa Xaa Xaa Xaa 90 Xaa | Xaa Xaa Xaa 75 Xaa Xaa | Xaa Xaa Xaa 60 Xaa Xaa Xaa | Xaa Xaa 45 Xaa Xaa Xaa | Xaa 30 Xaa Xaa Xaa Xaa 110 | 15 Xaa Xaa Xaa Xaa 95 Xaa | Xaa Xaa Xaa 80 Xaa Xaa |
| E> E> E> E> | 673 675 676 678 679 681 682 684 685 687 688 | 1 Glu Xaa Xaa Xaa 65 Xaa Xaa | Xaa Xaa 50 Xaa Xaa Xaa | Xaa Xaa 35 Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa Xaa 85 Xaa | Xaa Xaa Xaa 70 Xaa Xaa | Xaa Xaa 55 Xaa Xaa Xaa | Xaa Xaa 40 Xaa Xaa Xaa | Xaa 25 Xaa Xaa Xaa Xaa Xaa | 10 Xaa Xaa Xaa Xaa 90 Xaa | Xaa Xaa Xaa 75 Xaa Xaa | Xaa Xaa Xaa 60 Xaa Xaa Xaa | Xaa Xaa 45 Xaa Xaa Xaa | Xaa 30 Xaa Xaa Xaa Xaa 110 | 15 Xaa Xaa Xaa Xaa 95 Xaa | Xaa Xaa Xaa 80 Xaa Xaa |
| E> E> E> E> | 673 675 676 678 679 681 682 684 685 687 688 690 691 693 | 1 Glu Xaa Xaa Xaa 65 Xaa Xaa | Xaa Xaa 50 Xaa Xaa Xaa | Xaa 35 Xaa Xaa Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa 100 Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa 70 Xaa Xaa Xaa | Xaa Xaa 55 Xaa Xaa Xaa Xaa | Xaa 40 Xaa Xaa Xaa Xaa Xaa | Xaa 25 Xaa Xaa Xaa Xaa 105 Leu | 10 Xaa Xaa Xaa Xaa 90 Xaa Glu | Xaa Xaa Xaa 75 Xaa Xaa Glu | Xaa Xaa 60 Xaa Xaa Xaa Ser | Xaa 45 Xaa Xaa Xaa Xaa Gly 125 | Xaa 30 Xaa Xaa Xaa Xaa 110 Xaa | 15 Xaa Xaa Xaa Xaa 95 Xaa Xaa | Xaa Xaa Xaa 80 Xaa Xaa |
| E> E> E> E> | 673 675 676 678 679 681 682 684 685 687 688 690 691 693 | 1 Glu Xaa Xaa Xaa 65 Xaa Xaa | Xaa Xaa 50 Xaa Xaa Xaa | Xaa 35 Xaa Xaa Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa 100 Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa 70 Xaa Xaa Xaa | Xaa Xaa 55 Xaa Xaa Xaa Xaa | Xaa 40 Xaa Xaa Xaa Xaa Xaa | Xaa 25 Xaa Xaa Xaa Xaa 105 Leu | 10 Xaa Xaa Xaa Xaa 90 Xaa Glu | Xaa Xaa Xaa 75 Xaa Xaa Glu | Xaa Xaa 60 Xaa Xaa Xaa Ser | Xaa 45 Xaa Xaa Xaa Xaa Gly 125 | Xaa 30 Xaa Xaa Xaa Xaa 110 Xaa | 15 Xaa Xaa Xaa Xaa 95 Xaa Xaa | Xaa Xaa Xaa 80 Xaa Xaa |
| E> E> E> E> | 673 675 676 678 679 681 682 684 685 690 691 693 694 696 | 1 Glu Xaa Xaa Xaa 65 Xaa Xaa Xaa | Xaa Xaa 50 Xaa Xaa Xaa Xaa Xaa | Xaa 35 Xaa Xaa Xaa Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa 100 Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa 70 Xaa Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa 40 Xaa Xaa Xaa Xaa Xaa 120 Xaa | Xaa 25 Xaa Xaa Xaa Xaa 105 Leu | 10 Xaa Xaa Xaa Xaa 90 Xaa Glu Xaa | Xaa Xaa Xaa 75 Xaa Xaa Glu Xaa | Xaa Xaa 60 Xaa Xaa Xaa Xaa Xaa Ser | Xaa 45 Xaa Xaa Xaa Xaa Gly 125 Xaa | Xaa 30 Xaa Xaa Xaa Xaa 110 Xaa | Xaa Xaa Xaa Xaa Xaa 95 Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa |
| E> E> E> E> E> | 673 675 676 678 679 681 682 684 685 690 691 693 694 696 697 | 1 Glu Xaa Xaa Xaa 65 Xaa Xaa Xaa | Xaa Xaa 50 Xaa Xaa Xaa Xaa Xaa | Xaa 35 Xaa Xaa Xaa Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa 100 Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa 70 Xaa Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa 40 Xaa Xaa Xaa Xaa Xaa 120 Xaa | Xaa 25 Xaa Xaa Xaa Xaa 105 Leu | 10 Xaa Xaa Xaa Xaa 90 Xaa Glu Xaa | Xaa Xaa Xaa 75 Xaa Xaa Glu Xaa | Xaa Xaa 60 Xaa Xaa Xaa Xaa Xaa Ser | Xaa 45 Xaa Xaa Xaa Xaa Gly 125 Xaa | Xaa 30 Xaa Xaa Xaa Xaa 110 Xaa | Xaa Xaa Xaa Xaa Xaa 95 Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa |
| E> E> E> E> E> | 673 675 676 678 679 681 682 684 685 690 691 693 694 696 697 699 | Clu Xaa Xaa Xaa 65 Xaa Xaa Xaa Xaa Xaa | Xaa Xaa S0 Xaa Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa 100 Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa 70 Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa 40 Xaa Xaa Xaa Xaa 120 Xaa | Xaa 25 Xaa Xaa Xaa Xaa 105 Leu Xaa | 10 Xaa Xaa Xaa Xaa 90 Xaa Glu Xaa | Xaa Xaa Xaa 75 Xaa Xaa Glu Xaa Xaa 155 | Xaa Xaa 60 Xaa Xaa Xaa Ser Xaa 140 Xaa | Xaa 45 Xaa Xaa Xaa Xaa Gly 125 Xaa | Xaa 30 Xaa Xaa Xaa Xaa 110 Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Yaa Xaa Xaa | Xaa Xaa Xaa 80 Xaa Xaa Xaa Xaa Xaa Xaa Xaa |
| E> E> E> E> E> E> | 673 675 676 678 679 681 682 684 685 690 691 693 694 696 697 699 | Clu Xaa Xaa Xaa 65 Xaa Xaa Xaa Xaa Xaa | Xaa Xaa S0 Xaa Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa 100 Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa 70 Xaa Xaa Xaa Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa 40 Xaa Xaa Xaa Xaa 120 Xaa | Xaa 25 Xaa Xaa Xaa Xaa 105 Leu Xaa | 10 Xaa Xaa Xaa Xaa 90 Xaa Glu Xaa | Xaa Xaa Xaa 75 Xaa Xaa Glu Xaa Xaa 155 | Xaa Xaa 60 Xaa Xaa Xaa Ser Xaa 140 Xaa | Xaa 45 Xaa Xaa Xaa Xaa Gly 125 Xaa | Xaa 30 Xaa Xaa Xaa Xaa 110 Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Yaa Xaa Xaa | Xaa Xaa Xaa 80 Xaa Xaa Xaa Xaa Xaa Xaa Xaa |
| E> E> E> E> E> E> | 673 675 676 678 679 681 682 684 685 690 691 693 694 696 700 702 703 | 1 Glu Xaa Xaa Xaa 65 Xaa Xaa Xaa Xaa Xaa Xaa | Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa | Xaa 20 Xaa Xaa Xaa Xaa 100 Xaa Xaa Xaa | Xaa | Xaa Xaa Xaa 70 Xaa Xaa Xaa Xaa Xaa Xaa | Xaa | Xaa | Xaa 25 Xaa Xaa Xaa Xaa 105 Leu Xaa | 10 Xaa Xaa Xaa Xaa 90 Xaa Glu Xaa Xaa | Xaa Xaa Xaa 75 Xaa Xaa Glu Xaa Xaa 155 | Xaa Xaa 60 Xaa Xaa Xaa Ser Xaa 140 Xaa | Xaa 45 Xaa Xaa Xaa Xaa Gly 125 Xaa | Xaa 30 Xaa Xaa Xaa Xaa 110 Xaa Xaa | Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Clu | Xaa Xaa Xaa 80 Xaa Xaa Xaa Xaa Xaa Xaa Xaa |

some

Jane

RAW SEQUENCE LISTING

DATE: 04/24/2003

PATENT APPLICATION: US/09/830,502

TIME: 14:25:06

Input Set : A:\C26151.app
Output Set: N:\CRF4\04242003\I830502.raw

706

180

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/09/830,502

DATE: 04/24/2003 TIME: 14:25:07

Input Set : A:\C26151.app

Output Set: N:\CRF4\04242003\1830502.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of <220> to <223> is MANDATORY if n's or Xaa's are present. in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

```
Seq#:15; Xaa Pos. 18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36
Seq#:15; Xaa Pos. 37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55
Seq#:15; Xaa Pos. 56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74
Seq#:15; Xaa Pos. 75,76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93
Seq#:15; Xaa Pos. 94,95,96,97,98,99,100,101,102,103,104,105,106,107,108,109
Seq#:15; Xaa Pos. 110,111,112,113,114,115,116,117,118,119,120,126,127,128
Seq#:15; Xaa Pos. 129,130,131,132,133,134,135,136,137,138,139,140,141,142
Seq#:15; Xaa Pos. 143,144,145,146,147,148,149,150,151,152,153,154,155,156
Seq#:15; Xaa Pos. 157,158,159,160,161,162,163,164,165,166,167,168,169,170
Seq#:15; Xaa Pos. 171,172
Seq#:16; Xaa Pos. 18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36
Seq#:16; Xaa Pos. 37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55
Seq#:16; Xaa Pos. 56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74
Seq#:16; Xaa Pos. 75,76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93
Seq#:16; Xaa Pos. 94,95,96,97,98,99,100,101,102,103,104,105,106,107,108,109
Seq#:16; Xaa Pos. 110,111,112,113,114,115,116,117,118,119,120,129,130,131
Seq#:16; Xaa Pos. 132,133,134,135,136,137,138,139,140,141,142,143,144,145
Seq#:16; Xaa Pos. 146,147,148,149,150,151,152,153,154,155,156,157,158,159
Seq#:16; Xaa Pos. 160,161,162,163,164,165,166,167,168,169,170,171,172,173
Seq#:16; Xaa Pos. 174,175
Seq#:17; Xaa Pos. 18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36
Seq#:17; Xaa Pos. 37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55
Seq#:17; Xaa Pos. 56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74
Seq#:17; Xaa Pos. 75,76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93
Seq#:17; Xaa Pos. 94,95,96,97,98,99,100,101,102,103,104,105,106,107,108,109
Seq#:17; Xaa Pos. 110,111,112,113,114,115,116,117,118,119,120,126,127,128
Seq#:17; Xaa Pos. 129,130,131,132,133,134,135,136,137,138,139,140,141,142
Seq#:17; Xaa Pos. 143,144,145,146,147,148,149,150,151,152,153,154,155,156
Seq#:17; Xaa Pos. 157,158,159,160,161,162,163,164,165,166,167,168,169,170
Seq#:17; Xaa Pos. 171,172
Seq#:18; Xaa Pos. 18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36
Seq#:18; Xaa Pos. 37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55
Seq#:18; Xaa Pos. 56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74
Seq#:18; Xaa Pos. 75,76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93
Seq#:18; Xaa Pos. 94,95,96,97,98,99,100,101,102,103,104,105,106,107,108,109
Seq#:18; Xaa Pos. 110,111,112,113,114,115,116,117,118,119,120,126,127,128
Seq#:18; Xaa Pos. 129,130,131,132,133,134,135,136,137,138,139,140,141,142
Seq#:18; Xaa Pos. 143,144,145,146,147,148,149,150,151,152,153,154,155,156
Seq#:18; Xaa Pos. 157,158,159,160,161,162,163,164,165,166,167,168,169,170
Seg#:18; Xaa Pos. 171,172
Seq#:19; Xaa Pos. 18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36
Seq#:19; Xaa Pos. 37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55
Seq#:19; Xaa Pos. 56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74
```

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/09/830,502

DATE: 04/24/2003 TIME: 14:25:07

Input Set : A:\C26151.app

Output Set: N:\CRF4\04242003\I830502.raw

Seq#:19; Xaa Pos. 75,76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93
Seq#:19; Xaa Pos. 94,95,96,97,98,99,100,101,102,103,104,105,106,107,108,109
Seq#:19; Xaa Pos. 110,111,112,113,114,115,116,117,118,119,120,126,127,128
Seq#:19; Xaa Pos. 129,130,131,132,133,134,135,136,137,138,139,140,141,142
Seq#:19; Xaa Pos. 143,144,145,146,147,148,149,150,151,152,153,154,155,156
Seq#:19; Xaa Pos. 157,158,159,160,161,162,163,164,165,166,167,168,169,170
Seq#:19; Xaa Pos. 171,172
Seq#:20; Xaa Pos. 18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36

VERIFICATION SUMMARY DATE: 04/24/2003 TIME: 14:25:07 PATENT APPLICATION: US/09/830,502

Input Set : A:\C26151.app
Output Set: N:\CRF4\04242003\I830502.raw

| L:460 | M:340 E: | (46) "n" or | "Xaa" | used: | Feature | required, | for | SEQ | ID#:15 |
|-------|----------|-------------|-------|-------|---------|-----------|-----|-----|--------|
| M:340 | Repeated | in SeqNo=15 | | | | | | | |
| L:503 | M:340 E: | (46) "n" or | "Xaa" | used: | Feature | required, | for | SEQ | ID#:16 |
| M:340 | Repeated | in SeqNo=16 | | | | | | | • |
| L:546 | M:340 E: | (46) "n" or | "Xaa" | used: | Feature | required, | for | SEQ | ID#:17 |
| M:340 | Repeated | in SeqNo=17 | | | | | | | |
| L:589 | M:340 E: | (46) "n" or | "Xaa" | used: | Feature | required, | for | SEQ | ID#:18 |
| M:340 | Repeated | in SeqNo=18 | | | | | | | |
| L:632 | M:340 E: | (46) "n" or | "Xaa" | used: | Feature | required, | for | SEQ | ID#:19 |
| M:340 | Repeated | in SeqNo=19 | | | | | | | |
| L:675 | M:340 E: | (46) "n" or | "Xaa" | used: | Feature | required, | for | SEQ | ID#:20 |
| M:340 | Repeated | in SeqNo=20 | | | | | | | |